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Reviewer: Anne Corrigan

Timestamp: [year=2008; month=10; day=10; hr=12; min=42; sec=39; ms=857;  
]

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Application No: 10537839 Version No: 3.1

**Input Set:****Output Set:**

**Started:** 2008-10-10 12:40:22.231  
**Finished:** 2008-10-10 12:40:28.906  
**Elapsed:** 0 hr(s) 0 min(s) 6 sec(s) 675 ms  
**Total Warnings:** 36  
**Total Errors:** 0  
**No. of SeqIDs Defined:** 36  
**Actual SeqID Count:** 36

Error code	Error Description
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W 402	Undefined organism found in <213> in SEQ ID (2)
W 402	Undefined organism found in <213> in SEQ ID (3)
W 402	Undefined organism found in <213> in SEQ ID (4)
W 402	Undefined organism found in <213> in SEQ ID (5)
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W 402	Undefined organism found in <213> in SEQ ID (17)
W 402	Undefined organism found in <213> in SEQ ID (18)
W 402	Undefined organism found in <213> in SEQ ID (19)
W 402	Undefined organism found in <213> in SEQ ID (20)

**Input Set:**

**Output Set:**

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Error code	Error Description
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W 213	Artificial or Unknown found in <213> in SEQ ID (23)
W 213	Artificial or Unknown found in <213> in SEQ ID (24)
W 213	Artificial or Unknown found in <213> in SEQ ID (25)
W 213	Artificial or Unknown found in <213> in SEQ ID (26)
W 213	Artificial or Unknown found in <213> in SEQ ID (27)
W 213	Artificial or Unknown found in <213> in SEQ ID (28)
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W 213	Artificial or Unknown found in <213> in SEQ ID (30)
W 213	Artificial or Unknown found in <213> in SEQ ID (31)
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W 213	Artificial or Unknown found in <213> in SEQ ID (34)
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# SEQUENCE LISTING

<110> Hart, Derek Nigel John  
Kato, Masato

<120> DEC-205 (LY 75)/DCL-1 INTERGENIC SPLICE VARIANTS ASSOCIATED WITH HODGKIN'S  
DISEASE, AND USES THEREOF

<130> DAVI257.002APC

<140> 10537839

<141> 2006-05-18

<150> PCT/AU03/01634

<151> 2003-12-05

<150> AU2002953223

<151> 2002-12-06

<160> 36

<170> PatentIn version 3.1

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115 120 125			
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Val Trp Thr Tyr Ser Asp Thr Arg Cys Asp Ala Gly Trp Leu Pro Asn	
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370 375 380	
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Ala His Ala Lys Cys Lys Ala Phe Ser Ser Asp Leu Ile Ser Ile His	
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Ser Leu Ala Asp Val Glu Val Val Val Thr Lys Leu His Asn Glu Asp	
405 410 415	
atc aaa gaa gaa gtg tgg ata ggc ctt aag aac ata aac ata cca act	1296
Ile Lys Glu Glu Val Trp Ile Gly Leu Lys Asn Ile Asn Ile Pro Thr	
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435 440 445	
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450 455 460	
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465 470 475 480	
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acc tgt tac aag att tat gag gat gag gtc cct ttt gga aca aac tgc	1584
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625	630	635	640
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Asp Pro Cys Pro Glu Gly Trp Gln Ser Phe Pro Ala Ser Leu Ser Cys			
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agc cat gtg gat gaa ata aag gaa ttt ctt cac ttt tta acg gac cag			2112
Ser His Val Asp Glu Ile Lys Glu Phe Leu His Phe Leu Thr Asp Gln			
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Phe Ser Gly Gln His Trp Leu Trp Ile Gly Leu Asn Lys Arg Ser Pro			
705	710	715	720
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Asp Leu Gln Gly Ser Trp Gln Trp Ser Asp Arg Thr Pro Val Ser Thr			
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770 775 780	
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Tyr Glu Glu Ala Val Leu Tyr Cys Ala Ser Asn His Ser Phe Leu Ala	
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Thr Ile Thr Ser Phe Val Gly Leu Lys Ala Ile Lys Asn Lys Ile Ala	
850 855 860	
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Asn Ile Ser Gly Asp Gly Gln Lys Trp Trp Ile Arg Ile Ser Glu Trp	
865 870 875 880	
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Pro Ile Asp Asp His Phe Thr Tyr Ser Arg Tyr Pro Trp His Arg Phe	
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Ser Ala Ala Lys Val Gln Cys Ser Glu Gln Trp Ile Pro Phe Gln Asn	
945 950 955 960	
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Lys Cys Phe Leu Lys Ile Lys Pro Val Ser Leu Thr Phe Ser Gln Ala	
965 970 975	
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Ser Asp Thr Cys His Ser Tyr Gly Gly Thr Leu Pro Ser Val Leu Ser	



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Gly Phe Trp Lys Thr Val Asp Cys Asn Asp Asn Gln Pro Gly Ala  
1205 1210 1215

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